

Cryptococcus neoformans isolate A internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence

GenBank: MT466072.1

[FASTA Graphics](#)

[Go to:](#)

LOCUS MT466072 520 bp DNA linear PLN 19-MAY-2020
DEFINITION Cryptococcus neoformans isolate A internal transcribed spacer 1, partial sequence; 5.8S ribosomal RNA gene and internal transcribed spacer 2, complete sequence; and large subunit ribosomal RNA gene, partial sequence.

ACCESSION MT466072

VERSION MT466072.1

KEYWORDS .

SOURCE Cryptococcus neoformans (Filobasidiella neoformans var. bacillispora)

ORGANISM [Cryptococcus neoformans](#)

Eukaryota; Fungi; Dikarya; Basidiomycota; Agaricomycotina; Tremellomycetes; Tremellales; Cryptococcaceae; Cryptococcus; Cryptococcus neoformans species complex.

REFERENCE 1 (bases 1 to 520)

AUTHORS okolo,M.

TITLE Direct Submission

JOURNAL Submitted (14-MAY-2020) MEDICAL MICROBIOLOGY, JOS UNIVERSITY TEACHING HOSPITAL, LAMINGO, JOS, PLATEAU 930001, Nigeria

COMMENT ##Assembly-Data-START##

Sequencing Technology :: Sanger dideoxy sequencing

##Assembly-Data-END##

FEATURES Location/Qualifiers

source 1..520
/organism="Cryptococcus neoformans"
/mol_type="genomic DNA"
/isolate="A"
/db_xref="taxon:[5207](#)"

[misc RNA](#) <1..>520
/note="contains internal transcribed spacer 1, 5.8S ribosomal RNA, internal transcribed spacer 2, and large subunit ribosomal RNA"

ORIGIN

1 tgggacttcg ggtcmattta tctacccatc tacacctgtg aactgtttat gtgcttcggc
61 acgttttyacw caaacttcta aatgtaatga atgtaatctt attataacaa taataaaaact
121 ttcaacaacg gatctcttgg cttccacatc gatgaagaac gcagcgaaat gcgataagta
181 atgtgaattg cagaattcag tgaatcatcg aatctttgaa cgcaacttgc gccctttggg
241 attccgaagg gcatgcctgt ttgagagtca tgaaaatctc aatccctcgg gttttattac

```
301 ctgttggaact tggatttggg tgtttgccgc gacctgcaaa ggacgtcggc tcgccttaaa
361 tgtgttagtg ggaaggtgat tacctgtcag cccggcgtaa taagtttcgc tgggcctatg
421 gggtagtctt cggttgctg ataacaacca tctctttttg ttgacctca aatcaggtag
481 ggctaccgcg tgaacttaag catatcaata agcggaggaa
```

//